

# FIG. 1

CCCTCGACCTCGAGATCCATTGTGCTCTAAAGCAGAGTCTGGGTCTACCTACATATGGCACCCGAGGATA  
 CCTAGAGGCCCCATGCAAGAGAAAGGCCCTTGTGTTTCCAGGCACTGAGGACCGCAGTCCTTAATTCTCTGGC  
 AGTTCCTGAGATCTCAAGGAAAGCAGGTCAGCGAGGAGGCCCTGGGAGAGGAGGCATCCTACACCCAAT  
 CTTGTGGCCTGCTGCCCTAAGGAAACAGGAGACCA'GACAGCTATGC  
 TAACACTAGAAACCATGGCCAGTGAAGAAGAAATATGGGCCGAGGAACTGTGTGGTGTGGAGACCGGGC  
 CACAGGCTATCATTTCCACGCCCTGACTTGTGAGGGCTGCAAGGGCTTCTTCAGACGAACAGTCAGCAAA  
 ACCATTGGTCCCATCTGTCCGTTTGTGTGGAAGGTGAGGTCAAGCAAGGCCACAGAGACGCCACTGTCCAG  
 CCTGCAGGTTGCAGAAAGTGTCTAAATGTTGGCATGAGGAAAGACATGATACTGTCAAGCAGAAAGCCCTGGC  
 ATTGGCGCGAGCCAGACAGGCACAGCGCGGCGAGAGAAAGCATCTTTTGCAACTGAATCAGCAGCAGAAA  
 GAACTGGTCCAGATCCTCCTGGGGGCCACACTCGCCATGTGGGCCCATTTGTTTGACCAAGTTTGTGCAGT  
 TCAAGCCTCCAGCCTATCTGTTCATGTCATCACCGGCC'TTTCAGCCTCGGGCCCCGTTGTTCCTCTGCT  
 CACACACTTTGCAGATATCAACACGTTTATGTTGCAACAGATCATCAAGTTCAACCAAGGATCTGCCGCTC  
 TTCCGGTCCCTAACCATGGAGGACCAGATCCTCCTTCTCAAGGAGCGGCTGTGGAATATTGCATATCT  
 CACTCAACACTACGTTCTTCAAAACAGAGAAATTTCTTCTGTGGGCCCTCTTTGCTACAAGATGGAGGA  
 CGCAGTCCATGCAGGGTTCCAGTACGAGTTTGTGAGTCGATCCTCCACTTCCATAAAACCTGAAAGGA  
 CTGCATCTCCAGGAGCCTGAGTATGTGCTCATGGCTGCCACGGCCCTCTTCTCCCTGGTTTCTGTATGC  
 AAAGCTGATGGGCCCTGCTGGCTGACCTCCGGAGTATAAACAAATGCATACTCTATGAACCTTCAGCGCTTG  
 GAGGAACCTGTCTGCTATGACGCCCTGCTCGGGGAGATTTGCAGTTGAGGCCCAGGCTTGCAATCCTTTCC  
 CCAGACCCCAAGGAT'ACACTGGCCTGGAAAGGGTACAGCGCTGTGACCCCAACACAGCAGAAAGGAGCTTGG  
 GAGTGGCAATGAAATGCTGAACAGT (SEQ ID NO:1)

MTAMLTLTMASEEEYGPRNCVVCGDRATGYHFFHALTCEGCKGF  
 FRRTVSKTIGPICPFAGRCEVSKAQRRHCPACRLQKCLNVGMRKDM  
 ILSAEALALRRARQAQRRAEKASLQNNQQKELVQILLGAHTRHVG  
 PLFDQFVQFKPPAYLFMHHRPFQPRGPVLPLLTHFADINTFMVQQHK  
 FTKDLPLFRSLTMEDQISLLKGAAVEILHISLNTTFCLQTENFFCGPLC  
 YKMEDA VHAGFQYEFLESILHFHKNLKGHLHQEPEYVLMMAATA  
 LFSPGFQCMQS (SEQ ID NO:2)

FIG. 1

# Construct

GI Number(s): 2267577

Gene Family: Nuclear Hormone Receptor

Gene  
Subfamily: Orphan NHR

Gene Sequence: full-length cDNA, Mouse

underlined = deleted in targeting construct

[ ] = sequence flanking Neo insert in targeting construct

CCCTCGACCTCGAGATCCATTGTGCTCTAAAGCAGAGTCTGGGTCTACCTACATATGGC  
ACCGAGGATACCTAGAGGCCCATGCAAGAGAAGGCCCTTGTTTTCCAGGCACTGAGGAC  
CGCAGTCCCTAATTCCTGGCAGTTCCTGAGATCTCAAGGAAAGCAGGGTCAGCGAGGAGG  
CCTGGGGAGAGGAGGCATCTACACCCAATCTTGTGGCCTGCTGCCTAAGGGAAACAGCA  
GACCATGACAGCTATGCTAACACTAGAAACCATGGCCAGTGAAGAAGAATATGGGCCGAG  
GAACCTGTGTGGTGTGTGGAGACCGGGCCACAGGCTATCATTTCCACGCCCTGACTTGTGA  
GGGCTGCAAGGGCTTCTTCAGACGAACAGTCAGCAAAACCATTGGTCCCATCTGTCCGTT  
TGCTGGAAGGTGTGAGGTGAGCAAGGCCAGAGACGCCACTGTCCAGCCTGCAGGTTGCA  
GAAGTGTCTAAATGTTGGCATGAGGAAAGACATGATACTGTCAGCAGAAGCCCTGGCATT  
GCGGCGAGCCAGACAGGCACAGCGGGCCAGAGAAAGCATCTTTCAGCAATGAATCAGCA  
GCAGAAAGAACTGGTCCAGATCCTCCTGGGGGCCACACTCGCCATGTGGGCCCATTTGTT  
TGACCAGTTTGTGCAGTTCAAGCCTCCAGCCTATCTGTTTCATGCATCACCGGCCCTTTCCA  
GCCTCGGGGCCCCGTGTTGCCTCTGCTCACACACTTTGCAGATATCAACACGTTTATGGT  
GCAACAGATCATCAAGTTCACCAAGGATCTGCCGCTCTTCCGGTCCCTAACCATGGAGGA  
CCAGATCTCCCTTCTCAAGGGAGCGGCTGTGGAATATTCATATCTCACTCAACACTAC  
GTTCTGTCTTCAACAGAGAATTTCTTCTGTGGGCCTCTTTCAGTACAAGATGGAGGACGC  
AGTCCATGCAGGGTTCCAGTACGAGTTTTTGGAGTCGATCCTCCACTTCCATAAAAAACCT  
GAAAGGACTGCATCTCCAGGAGCCTGAGTATGTGCTCATGGCTGCCACGGCCCTCTTCTC  
CCCTGGTTTCTGTATGCAAGCTGATGGGCCTGCTGGCTGACCTCCGGAGTATAAACAAT  
GCATACTCCTATGAACCTTCAGCGCTTGGAGGAACGTCTGCTATGACGCCGCTGCTCGGG  
GAGATTTGCAGTTGAGGCCAGGCTTGCATCCTTTCCCCAGACCCCCAGGGATACACTGG  
CCTGGAAAGGGTACAGCGCTGGACCCACACAGCAGAAGGAGCTTGGGAGTGGCAATGAA  
ATGCTGAACAGT

Gene Sequence  
Structure \*

282 bp

Sequence Deleted

403 bp

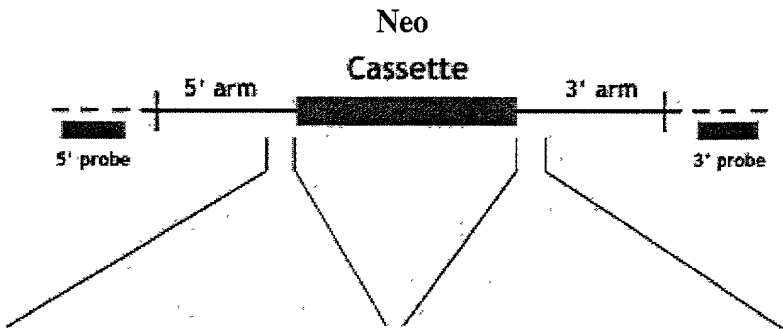
Size of full-length  
cDNA: 1332 bp

FIG. 2A

Targeting Vector\* (genomic sequence)

Construct Number: 138

Arm Length:  
5': 3.5 kb  
3': 1.9 kb



———— Targeting Vector  
----- Endogenous Locus

\* Not drawn to scale

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5'>TTGGAGGCCAGAGACAAAAAG
CAACATTTTGTCTTTAATGTCCT
CAGTGCTGGGGAGCCCCGGTGCAG
GCTGGGCAGTCTTGGAAGAGATT
CTGTAGAGGAGAGAGAAGAGAGTC
CTATGGCCCAGTGCTGATTCTCAA
CTCCTCCCACATTCAGGAGACCAT
GACAGCTATGCTAACA:TAGAAAC
CATGGCCAGTG<3'
(SEQ ID NO. 3)
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5'>GGTCCCATCTGTCCGTTTGCT
GGAAGGTGTGAGGTCAGCAAGGCC
CAGAGACGCCACTGTCCAGCCTGC
AGGTTGCAGAAGTGTCTAAATGTT
GGCATGAGGAAAGACAGTGAGTTG
GTCCCTACCGTTCCAGGAGTCCAT
TGGTGCTTCTCTGCTTTTGTTCAG
GTTCCACAGCAGGCAGAGTGAGC
TGTTGGAAACC<3'
(SEQ ID NO. 4)
```

FIG. 2B

# necropsy - spleen weight/body weight

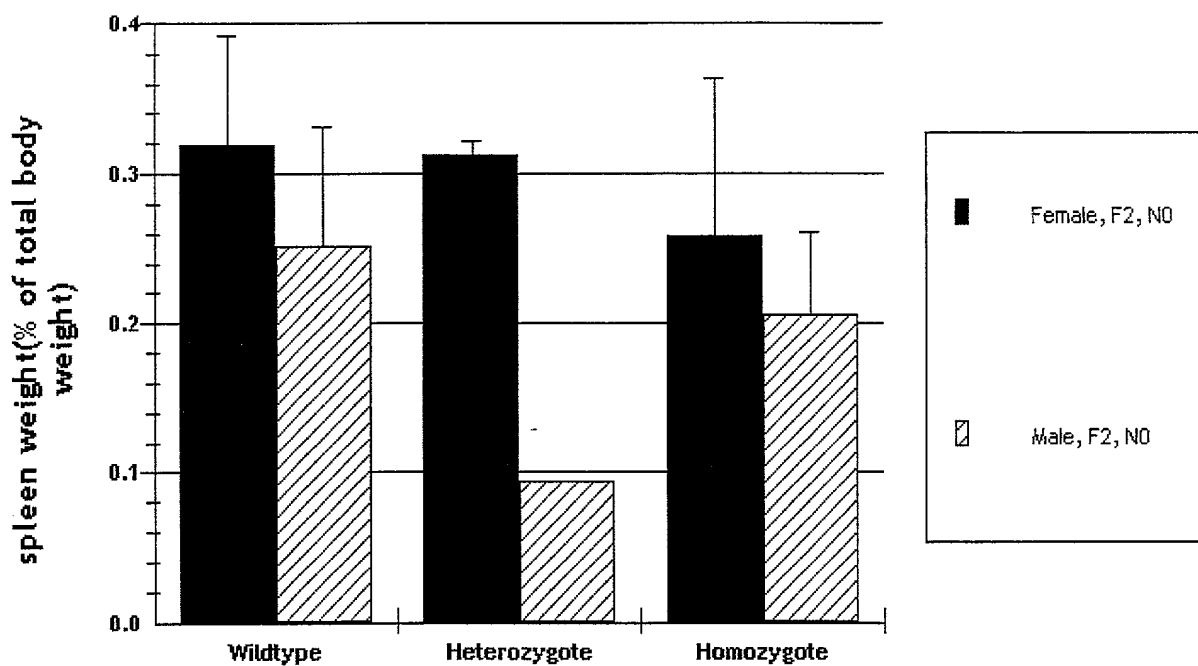


FIG. 3

# necropsy - thymus weight/body weight

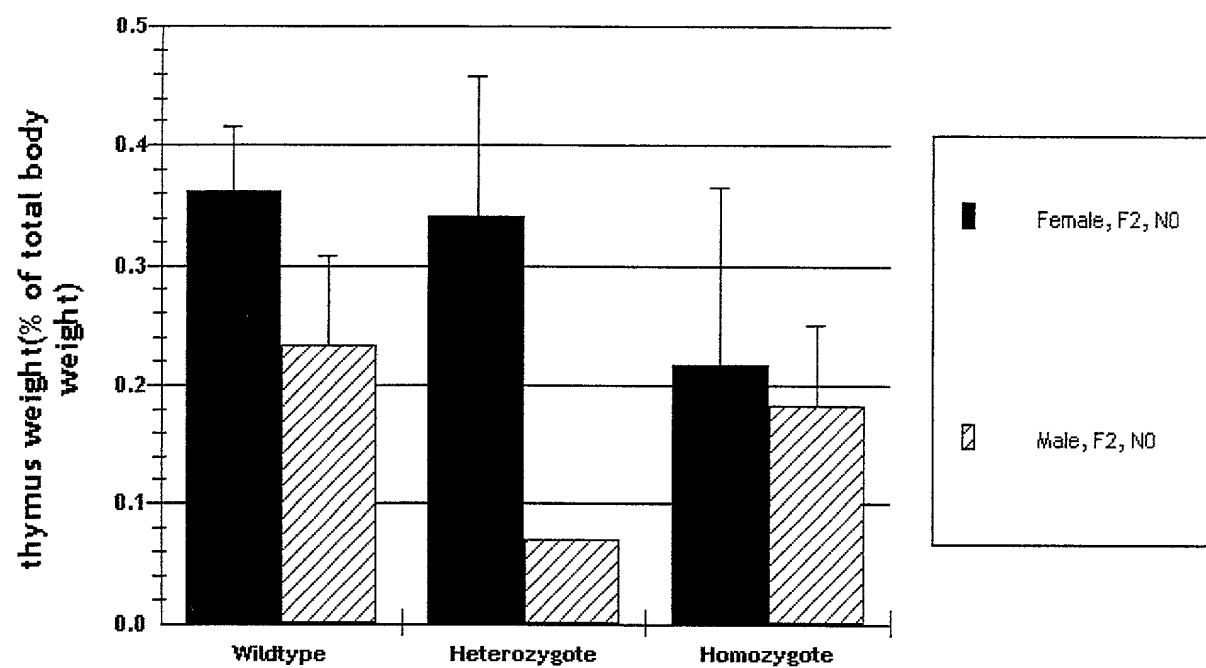


FIG. 4